SEQUENCE LISTING

<110> HITOSHI, KIYOI NAOE, TOMOKI TOWATARI, MASAYUKI KITAMURA, TOSHIO										I	AYUK:	OKI MAS	TOMARI,	AOE, TAWC	NZ T(<110
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tca gat aat gag tac ttc tac gtt gat ttc aga gaa tat gaa tat gat 96 Ser Asp Asn Glu Tyr Phe Tyr Val Asp Phe Arg Glu Tyr Glu Tyr Asp 20 25 30			Glu					Asp					Glu			
ctc aaa tgg gag ttt cca aga gaa aat tgc tcc tca gat aat gag tac 14 Leu Lys Trp Glu Phe Pro Arg Glu Asn Cys Ser Ser Asp Asn Glu Tyr 35 40 45				Asp					Glu					Trp		
ttc tac gtt gat ttc aga gaa tat gaa tat gat ctc aaa tgg gag ttt 19 Phe Tyr Val Asp Phe Arg Glu Tyr Glu Tyr Asp Leu Lys Trp Glu Phe 50 55 60					Leu					Glu					Tyr	
cca aga gaa aat tta gag ttt ggg aag gta cta gga tca ggt gct ttt 24 Pro Arg Glu Asn Leu Glu Phe Gly Lys Val Leu Gly Ser Gly Ala Phe 65 70 75 80	Phe					Leu					Glu					Pro

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gga aaa gtg atg aac gca aca gct tat gga att agc aaa aca gga gtc
Gly Lys Val Met Asn Ala Thr Ala Tyr Gly Ile Ser Lys Thr Gly Val
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Ser Ile Gln Val Ala Val Lys Met Leu Lys
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Leu Lys Trp Glu Phe Pro Arg Glu Asn Cys Ser Ser Asp Asn Glu Tyr
Phe Tyr Val Asp Phe Arg Glu Tyr Glu Tyr Asp Leu Lys Trp Glu Phe
                         55
Pro Arg Glu Asn Leu Glu Phe Gly Lys Val Leu Gly Ser Gly Ala Phe
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ctc aaa agc Leu Lys Ser 35		-					Phe	_	_	144
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Ser Asp Asn Glu Tyr Phe Tyr Val Asp Phe Arg Glu Tyr Glu Met Gly
atg ggg gga gaa tgt aat ccc ggg aga caa gat ctc aaa tgg gag ttt
                                                                   144
Met Gly Glu Cys Asn Pro Gly Arg Gln Asp Leu Lys Trp Glu Phe
cca aga gaa aat tta gag ttt ggg aag gta cta gga tca ggt gct ttt
                                                                   192
Pro Arg Glu Asn Leu Glu Phe Gly Lys Val Leu Gly Ser Gly Ala Phe
gga aaa gtg atg aac gca aca gct tat gga att agc aaa aca gga gtc
                                                                   240
Gly Lys Val Met Asn Ala Thr Ala Tyr Gly Ile Ser Lys Thr Gly Val
tca atc cag gtt gcc gtc aaa atg ctg aaa g
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                                                                   96
Ser Asp Asn Glu Tyr Phe Tyr Val Asp Phe Arg Glu Tyr Asp Glu Tyr
ttc tac gtt gat ttc aga gaa tat gaa tat gat ctc aaa tgg gag ttt
                                                                   144
Phe Tyr Val Asp Phe Arg Glu Tyr Glu Tyr Asp Leu Lys Trp Glu Phe
         35
cca aga gaa aat tta gag ttt ggg aag gta cta gga tca ggt gct ttt
                                                                   192
Pro Arg Glu Asn Leu Glu Phe Gly Lys Val Leu Gly Ser Gly Ala Phe
gga aaa gtg atg aac gca aca gct tat gga att agc aaa aca gga gtc
                                                                   240
Gly Lys Val Met Asn Ala Thr Ala Tyr Gly Ile Ser Lys Thr Gly Val
 65
                     70
                                          75
tca atc cag gtt gcc gtc aaa atg ctg aaa g
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Ser Ile Gln Val Ala Val Lys Met Leu Lys
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Phe Tyr Val Asp Phe Arg Glu Tyr Glu Tyr Asp Leu Lys Trp Glu Phe 40 Pro Arg Glu Asn Leu Glu Phe Gly Lys Val Leu Gly Ser Gly Ala Phe Gly Lys Val Met Asn Ala Thr Ala Tyr Gly Ile Ser Lys Thr Gly Val 70 Ser Ile Gln Val Ala Val Lys Met Leu Lys 85 <210> 9 <211> 25 <212> DNA <213> Artificial Sequence <223> Description of Artificial Sequence: A primer for amplifying human FLT3/ITD genes caacaattgg tgtttgtctc ctctt 25 <210> 10 <211> 25 <212> DNA <213> Artificial Sequence <220>

<223> Description of Artificial Sequence: A primer for

amplifying human FLT3/ITD genes.

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